



SEQUENCE LISTING

<110> Bulla, Lee
Candas, Mehmet

<120> Pectinophora gossypiella (Pink Bollworm) Bacillus
thuringiensis toxin receptor BT-R2

<130> 52418-20003.00

<140> US 09/696,115

<141> 2000-10-24

<150> US 60/161,564

<151> 1999-10-26

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<170> PatentIn version 3.0

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<212> DNA

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Lys	Ile	Phe	Tyr	Ile	Gln	Gly	Ala	Asn	Ile	Pro	Gly	Glu	Trp	Ile	Arg
225					230					235					240
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Leu	Pro	Arg	Gly	Ser	Pro	Asn	Val	Glu	Glu	Asn	Val	Pro	Glu	Gly
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Leu	Arg	Phe	Glu	Ile	Asp	Trp	Thr	Thr	Ser	Tyr	Ala	Thr	Lys	Gln
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Arg	Arg	Tyr	Asn	Leu	Tyr	Tyr	Thr	Val	Val	Ala	Thr	Asp	Arg	Cys
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Ala	Glu	Asp	Pro	Asp	Asp	Cys	Pro	Asp	Asp	Pro	Thr	Tyr	Trp	Glu
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Arg	Asp	Asp	Ile	Tyr	His	Thr	Ile	Arg	Tyr	Gln	Ile	Asn	Tyr	Ala	Val	915	920	925
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		500						505					510		
Glu	Gln	Pro	Ile	Phe	Glu	His	Ala	Val	Gln	Thr	Val	Thr	Phe	Asp	Glu
		515					520					525			
Thr	Glu	Gly	Glu	Gly	Phe	Phe	Val	Ala	Lys	Ala	Val	Ala	His	Asp	Arg
	530					535					540				
Asp	Ile	Gly	Asp	Val	Val	Glu	His	Thr	Leu	Leu	Gly	Asn	Ala	Val	Asn

545	Phe	Leu	Thr	Ile	Asp	550	Lys	Leu	Thr	Gly	Asp	555	Ile	Arg	Val	Ser	Ala	Asn	560
					565						570							575	
	Asp	Ser	Phe	Asn	Tyr		His	Arg	Glu	Ser	Glu		Leu	Phe	Val	Gln	Val	Arg	
				580						585								590	
	Ala	Thr	Asp	Thr	Leu		Gly	Glu	Pro	Phe	His		Thr	Ala	Thr	Ser	Gln	Leu	
			595						600							605			
	Val	Ile	Arg	Leu	Asn		Asp	Ile	Asn	Asn	Thr		Pro	Pro	Thr	Leu	Arg	Leu	
		610						615							620				
	Pro	Arg	Gly	Ser	Pro		Gln	Val	Glu	Glu	Asn		Val	Pro	Asp	Gly	His	Val	
		625						630							635			640	
	Ile	Thr	Gln	Glu	Leu		Arg	Ala	Thr	Asp	Pro		Asp	Thr	Thr	Ala	Asp	Leu	
					645						650							655	
	Arg	Phe	Glu	Ile	Asn		Trp	Asp	Thr	Ser	Phe		Ala	Thr	Lys	Gln	Gly	Arg	
				660						665								670	
	Gln	Ala	Asn	Pro	Asp		Glu	Phe	Arg	Asn	Cys		Val	Glu	Ile	Glu	Thr	Ile	
			675						680							685			
	Phe	Pro	Glu	Ile	Asn		Asn	Arg	Gly	Leu	Ala		Ile	Gly	Arg	Val	Val	Ala	
		690						695							700				
	Arg	Glu	Ile	Arg	His		Asn	Val	Thr	Ile	Asp		Tyr	Glu	Glu	Phe	Glu	Val	
		705						710						715				720	
	Leu	Ser	Leu	Thr	Val		Arg	Val	Arg	Asp	Leu		Asn	Thr	Val	Tyr	Gly	Asp	
					725						730							735	
	Asp	Tyr	Asp	Glu	Ser		Met	Leu	Thr	Ile	Thr		Ile	Ile	Asp	Met	Asn	Asp	
				740						745								750	
	Asn	Ala	Pro	Val	Trp		Val	Glu	Gly	Thr	Leu		Glu	Gln	Asn	Phe	Arg	Val	
			755						760									765	
	Arg	Glu	Met	Ser	Ala		Gly	Gly	Leu	Val	Val		Gly	Ser	Val	Arg	Ala	Asp	
		770						775							780				
	Asp	Ile	Asp	Gly	Pro		Leu	Tyr	Asn	Gln	Val		Arg	Tyr	Thr	Ile	Phe	Pro	
		785						790						795				800	
	Arg	Glu	Asp	Thr	Asp		Lys	Asp	Leu	Ile	Met		Ile	Asp	Phe	Leu	Thr	Gly	
					805						810							815	
	Gln	Ile	Ser	Val	Asn		Thr	Ser	Gly	Ala	Ile		Asp	Ala	Asp	Thr	Pro	Pro	
				820						825								830	
	Arg	Phe	His	Leu	Tyr		Tyr	Thr	Val	Val	Ala		Ser	Asp	Arg	Cys	Ser	Thr	
			835						840									845	
	Glu	Asp	Pro	Ala	Asp		Cys	Pro	Pro	Asp	Pro		Thr	Tyr	Trp	Glu	Thr	Glu	
		850						855							860				
	Gly	Asn	Ile	Thr	Ile		His	Ile	Thr	Asp	Thr		Asn	Asn	Lys	Val	Pro	Gln	
		865						870						875				880	
	Ala	Glu	Thr	Thr	Lys		Phe	Asp	Thr	Val	Val		Tyr	Ile	Tyr	Glu	Asn	Ala	
					885						890							895	
	Thr	His	Leu	Asp	Glu		Val	Val	Thr	Leu	Ile		Ala	Ser	Asp	Leu	Asp	Arg	
				900						905								910	
	Asp	Glu	Ile	Tyr	His		Thr	Val	Ser	Tyr	Val		Ile	Ile	Asn	Tyr	Ala	Val	
			915						920							925			
	Asn	Pro	Arg	Leu	Met		Asn	Phe	Phe	Ser	Val		Asn	Arg	Glu	Thr	Gly	Leu	
		930						935								940			
	Val	Tyr	Val	Asp	Tyr		Glu	Thr	Gln	Gly	Ser		Gly	Glu	Val	Leu	Asp	Arg	
		945						950						955				960	
	Asp	Gly	Asp	Glu	Pro		Thr	His	Arg	Ile	Phe		Phe	Asn	Leu	Ile	Asp	Asn	
					965						970							975	
	Phe	Met	Gly	Glu	Gly		Glu	Gly	Asn	Arg	Asn		Gln	Asn	Asp	Thr	Glu	Val	
			980						985									990	
	Leu	Val	Ile	Leu	Leu		Asp	Val	Asn	Asp	Asn		Ala	Pro	Glu	Leu	Pro	Pro	
			995						1000							1005			
	Pro	Ser	Glu	Leu	Ser		Trp	Thr	Ile	Ser	Glu		Asn	Leu	Lys	Gln	Gly	Val	
		1010						1015								1020			
	Arg	Leu	Glu	Pro	His		Ile	Phe	Ala	Pro	Asp		Arg	Asp	Glu	Pro	Asp	Thr	
		1025						1030						1035				1040	

Asp Asn Ser Arg Val Gly Tyr Glu Ile Leu Asn Leu Ser Thr Glu Arg
 1045 1050 1055
 Asp Ile Glu Val Pro Glu Leu Phe Val Met Ile Gln Ile Ala Asn Val
 1060 1065 1070
 Thr Gly Glu Leu Glu Thr Ala Met Asp Leu Lys Gly Tyr Trp Gly Thr
 1075 1080 1085
 Tyr Ala Ile His Ile Arg Ala Phe Asp His Gly Ile Pro Gln Met Ser
 1090 1095 1100
 Met Asn Glu Thr Tyr Glu Leu Ile Ile His Pro Phe Asn Tyr Tyr Ala
 1105 1110 1115 1120
 Pro Glu Phe Val Phe Pro Thr Asn Asp Ala Val Ile Arg Leu Ala Arg
 1125 1130 1135
 Glu Arg Ala Val Ile Asn Gly Val Leu Ala Thr Val Asn Gly Glu Phe
 1140 1145 1150
 Leu Glu Arg Ile Ser Ala Thr Asp Pro Asp Gly Leu His Ala Gly Val
 1155 1160 1165
 Val Thr Phe Gln Val Val Gly Asp Glu Glu Ser Gln Arg Tyr Phe Gln
 1170 1175 1180
 Val Val Asn Asp Gly Glu Asn Leu Gly Ser Leu Arg Leu Leu Gln Ala
 1185 1190 1195 1200
 Val Pro Glu Glu Ile Arg Glu Phe Arg Ile Thr Ile Arg Ala Thr Asp
 1205 1210 1215
 Gln Gly Thr Asp Pro Gly Pro Leu Ser Thr Asp Met Thr Phe Arg Val
 1220 1225 1230
 Val Phe Val Pro Thr Gln Gly Glu Pro Arg Phe Ala Ser Ser Glu His
 1235 1240 1245
 Ala Val Ala Phe Ile Glu Lys Ser Ala Gly Met Glu Glu Ser His Gln
 1250 1255 1260
 Leu Pro Leu Ala Gln Asp Ile Lys Asn His Leu Cys Glu Asp Asp Cys
 1265 1270 1275 1280
 His Ser Ile Tyr Tyr Arg Ile Ile Asp Gly Asn Ser Glu Gly His Phe
 1285 1290 1295
 Gly Leu Asp Pro Val Arg Asn Arg Leu Phe Leu Lys Lys Glu Leu Ile
 1300 1305 1310
 Arg Glu Gln Ser Ala Ser His Thr Leu Gln Val Ala Ala Ser Asn Ser
 1315 1320 1325
 Pro Asp Gly Gly Ile Pro Leu Pro Ala Ser Ile Leu Thr Val Thr Val
 1330 1335 1340
 Thr Val Arg Glu Ala Asp Pro Arg Pro Val Phe Val Arg Glu Leu Tyr
 1345 1350 1355 1360
 Thr Ala Gly Ile Ser Thr Ala Asp Ser Ile Gly Arg Glu Leu Leu Arg
 1365 1370 1375
 Leu His Ala Thr Gln Ser Glu Gly Ser Ala Ile Thr Tyr Ala Ile Asp
 1380 1385 1390
 Tyr Asp Thr Met Val Val Asp Pro Ser Leu Glu Ala Val Arg Gln Ser
 1395 1400 1405
 Ala Phe Val Leu Asn Ala Gln Thr Gly Val Leu Thr Leu Asn Ile Gln
 1410 1415 1420
 Pro Thr Ala Thr Met His Gly Leu Phe Lys Phe Glu Val Thr Ala Thr
 1425 1430 1435 1440
 Asp Thr Ala Gly Ala Gln Asp Arg Thr Asp Val Thr Val Tyr Val Val
 1445 1450 1455
 Ser Ser Gln Asn Arg Val Tyr Phe Val Phe Val Asn Thr Leu Gln Gln
 1460 1465 1470
 Val Glu Asp Asn Arg Asp Phe Ile Ala Asp Thr Phe Ser Ala Gly Phe
 1475 1480 1485
 Asn Met Thr Cys Asn Ile Asp Gln Val Val Pro Ala Asn Asp Pro Val
 1490 1495 1500
 Thr Gly Val Ala Leu Glu His Ser Thr Gln Met Arg Gly His Phe Ile
 1505 1510 1515 1520
 Arg Asp Asn Val Pro Val Leu Ala Asp Glu Ile Glu Gln Ile Arg Ser

	1525		1530		1535
Asp Leu Val Leu Leu Ser Ser Ile Gln Thr Thr Leu Ala Ala Arg Ser					
	1540		1545		1550
Leu Val Leu Asp Leu Leu Thr Asn Ser Ser Pro Asp Ser Ala Pro Asp					
	1555		1560		1565
Ser Ser Leu Thr Val Tyr Val Leu Ala Ser Leu Ser Ala Val Leu Gly					
	1570		1575		1580
Phe Met Cys Leu Val Leu Leu Leu Thr Phe Ile Ile Arg Thr Arg Ala					
1585		1590		1595	1600
Leu Asn Arg Arg Leu Glu Ala Leu Ser Met Thr Lys Tyr Gly Ser Leu					
	1605		1610		1615
Asp Ser Gly Leu Asn Arg Ala Gly Ile Ala Ala Pro Gly Thr Asn Lys					
	1620		1625		1630
His Thr Val Glu Gly Ser Asn Pro Ile Phe Asn Glu Ala Ile Lys Thr					
	1635		1640		1645
Pro Asp Leu Asp Ala Ile Ser Glu Gly Ser Asn Asp Ser Asp Leu Ile					
	1650		1655		1660
Gly Ile Glu Asp Leu Pro His Phe Gly Asn Val Phe Met Asp Pro Glu					
1665		1670		1675	1680
Val Asn Glu Lys Ala Asn Gly Tyr Pro Glu Val Ala Asn His Asn Asn					
	1685		1690		1695
Asn Phe Ala Phe Asn Pro Thr Pro Phe Ser Pro Glu Phe Val Asn Gly					
	1700		1705		1710
Gln Phe Arg Lys Ile					
	1715				